

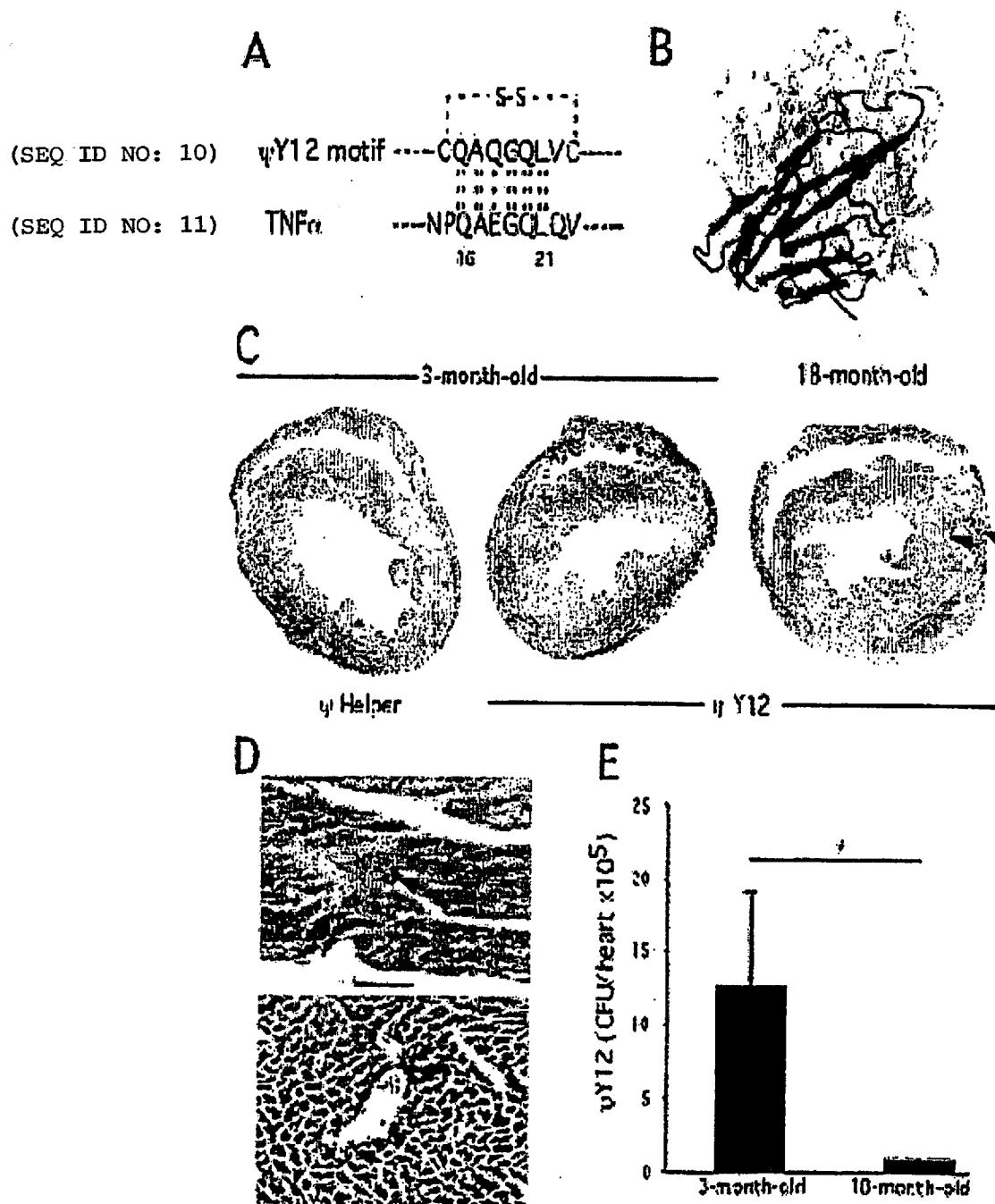
## Figure 1

1 RTPSDKPVAH VVANPQAEGQ LQWLNRRANA LLANGVELRD NQLVVVPSEG  
51 YLIYSQVLFK GQGCPSTHVL LTHTISRIAV SYQTKVNLLS AIKSPCQRET  
101 PEGAEAAKPWF EPIYILGGVFQ LEKGDRILSAE INRPDYLLFA ESGQQVYFGII  
151 AL (SEQ ID NO: 6)

## Figure 2

1 HSDPARRGQL SVCDSISEWV TAADKKKTAVD MSGGGTVTVLE KVPVSKGQLK  
51 QYFYETKCNP MGYTKEGCRG IDKRWNSQC RTTQSYVRAL TMDSKKRIGW  
101 RFIRIDTSCV CTLTIKRGR (SEQ ID NO: 7)

## FIGURE 3



## FIGURE 6

